Please amend the application as follows:

IN THE CLAIMS:

Please cancel claims 56-63, without prejudice or disclaimer and add the following claims:

--64. A modular replicable vector, comprising (a) a plurality of vector modules, wherein each vector module is flanked by DNA cleavage sites unique within said vector, and (b) DNA cleavage sites for insertion of a polypeptide-encoding nucleotide sequence wherein said DNA cleavage sites are unique within said vector.

65. A modular replicable vector adapted for insertion of a nucleotide sequence encoding a (poly)peptide, comprising; (a) a plurality of vector modules, wherein each vector module is flanked by DNA cleavage sites unique within said vector and said nucleotide sequence; and (b) DNA cleavage sites for insertion of said nucleotide sequence encoding a (poly)peptide, wherein said DNA cleavage sites are unique within said vector.

66. A modular replicable vector comprising: (a) a nucleotide sequence encoding a (poly)peptide, wherein said nucleotide sequence is flanked by DNA cleavage sites unique within said vector and said nucleotide sequence; and (b) a plurality of vector modules, wherein each vector module is flanked by DNA cleavage sites unique within said vector and said nucleotide sequence.

67. The vector according to claim 64, comprising (a) an origin of replication selected from the group consisting of an origin of double-stranded replication for a low copy number plasmid and an origin of double-stranded replication for a high copy number plasmid; and (b) a plurality of vector modules selected from the group consisting of a promoter element, an operator element, a repressor element, a terminator element, a resistance gene, a recombination site, a filamentous phage gene III, a truncated filamentous phage gene III, a signal sequence, a purification tag, a detection tag, and a sequence encoding an additional (poly)peptide moiety.

68. The vector according to claim 67, wherein said additional moiety is selected from the group consisting of a toxin, a cytokine, a reporter enzyme, a metal-binding moiety, a peptide, a tag suitable for detection and/or purification, a homo-association domain and a hetero-association domain.

69. The vector according to claim 64, wherein said vector is pMCS (SEQ ID NO: 264).

70. The vector according to claim 64, wherein said vector is selected from the group consisting of pCAL 4 (SEQ ID NO: 274), pCALO-1 (SEQ ID NO: 294), pCALO-2 (SEQ ID NO: 296), and pCALO-3 (SEQ ID NO: 299).

71. The vector according to claim 65, wherein said nucleotide sequence encoding a (poly)peptide encodes a (poly)peptide selected from the group consisting of an immunoglobulin light chain variable region, an immunoglobulin heavy chain variable region, and an immunoglobulin constant chain region.

72. The vector according to claim 71, wherein said nucleotide sequence encoding a (poly)peptide encodes an immunoglobulin light chain variable region comprising a modular sequence of four consensus framework regions interspaced by three complementary determining regions CDR1, CDR2, and CDR3,
wherein said nucleotide sequence comprises DNA cleavage sites at the boundary between each consensus framework region and each complementary determining region, and wherein each of said cleavage sites is unique within said vector.